

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/807,660

Source: PCT09

Date Processed by STIC: 5-1-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/807,660

DATE: 05/01/2001

TIME: 11:37:50

Input Set : A:\23,495 PCT.txt

Output Set: N:\CRF3\05012001\I807660.raw

Does Not Comply
Corrected Diskette Needed
see pp. 1, 2

```

3 <110> APPLICANT: Maliszewski, Charles R.
4      Gayle III, Richard B.
5      Marcus, Aaron J.
6      Immunex Corporation
7      Cornell Research Foundation, Inc.
9 <120> TITLE OF INVENTION: Methods of Inhibiting Platelet Activation and
10      Recruitment
12 <130> FILE REFERENCE: 23,495 PCT
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/807,660
C--> 15 <141> CURRENT FILING DATE: 2001-04-16
17 <150> PRIOR APPLICATION NUMBER: US 60/104,585
18 <151> PRIOR FILING DATE: 1998-10-16
20 <150> PRIOR APPLICATION NUMBER: US 60/107,466
21 <151> PRIOR FILING DATE: 1998-11-06
23 <150> PRIOR APPLICATION NUMBER: US 60/149,010
24 <151> PRIOR FILING DATE: 1999-08-13
26 <160> NUMBER OF SEQ ID NOS: 31
28 <170> SOFTWARE: PatentIn Ver. 2.0

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ERRORED SEQUENCES

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1925 <210> SEQ ID NO: 31
1926 <211> LENGTH: 58
1927 <212> TYPE: PRT
1928 <213> ORGANISM: Homo sapiens
1930 <400> SEQUENCE: 31
1931 Met Ala Thr Ser Trp Gly Thr Val Phe Phe Met Leu Val Val Ser Cys
1932 1      5      10      15
1934 Val Cys Ser Ala Val Ser His Arg Asn Gln Gln Thr Trp Phe Glu Gly
1935      20      25      30
1937 Ile Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Ser Thr Leu
1938      35      40      45
1940 Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr
1941      50      55
E--> 1945 1
E--> 1948 1

```

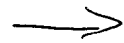
→ Delete extraneous numerals at the end of the file. They are causing an invalid amino acid count for sequence #31.

<210> 6 Seg #6 (Partially shown)

<211> 454

<212> PRT

<213> Artificial Sequence



missing mandatory <220>, <223>
feature to explain the source

<400> 6

Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln Leu Thr Ser Ser Thr
1 5 10 15

of the artificial

Gln Asn Lys Ala Leu Pro Glu Asn Val Lys Tyr Gly Ile Val Leu Asp
20 25 30

sequence.



Ala Gly Ser Ser His Thr Ser Leu Tyr Ile Tyr Lys Trp Pro Ala Glu
35 40 45

Note: Error

Lys Glu Asn Asp Thr Gly Val Val His Gln Val Glu Glu Cys Arg Val
50 55 60

repeated in
Sequence #5
8, 26

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/807,660

DATE: 05/01/2001

TIME: 11:37:52

Input Set : A:\23,495 PCT.txt

Output Set: N:\CRF3\05012001\I807660.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:620 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:620 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:850 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:850 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:1426 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1426 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:1945 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:31
M:332 Repeated in SeqNo=31